

This Page Is Inserted by IFW Operations
and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

**As rescanning documents *will not* correct images,
please do not report the images to the
Image Problem Mailbox.**

THIS PAGE BLANK (USPTO)

GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: June 18, 2003, 03:16:37 ; Search time 33.4567 Seconds
(without alignments)
1215.770 Million cell updates/sec

Title: US-09-807-933B-5
Perfect score: 1956
Sequence: 1 MKFLTASXALALAVGTEM.....TYKQVCPKAITAKSGCSRK 360

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database :

1: A_Geneseq_101002.*
2: /SID2/gcgdata/geneseq/geneseq-emb1/AA1980.DAT.*
3: /SID2/gcgdata/geneseq/geneseq-emb1/AA1981.DAT.*
4: /SID2/gcgdata/geneseq/geneseq-emb1/AA1982.DAT.*
5: /SID2/gcgdata/geneseq/geneseq-emb1/AA1983.DAT.*
6: /SID2/gcgdata/geneseq/geneseq-emb1/AA1984.DAT.*
7: /SID2/gcgdata/geneseq/geneseq-emb1/AA1985.DAT.*
8: /SID2/gcgdata/geneseq/geneseq-emb1/AA1986.DAT.*
9: /SID2/gcgdata/geneseq/geneseq-emb1/AA1987.DAT.*
10: /SID2/gcgdata/geneseq/geneseq-emb1/AA1988.DAT.*
11: /SID2/gcgdata/geneseq/geneseq-emb1/AA1989.DAT.*
12: /SID2/gcgdata/geneseq/geneseq-emb1/AA1990.DAT.*
13: /SID2/gcgdata/geneseq/geneseq-emb1/AA1991.DAT.*
14: /SID2/gcgdata/geneseq/geneseq-emb1/AA1992.DAT.*
15: /SID2/gcgdata/geneseq/geneseq-emb1/AA1993.DAT.*
16: /SID2/gcgdata/geneseq/geneseq-emb1/AA1994.DAT.*
17: /SID2/gcgdata/geneseq/geneseq-emb1/AA1995.DAT.*
18: /SID2/gcgdata/geneseq/geneseq-emb1/AA1996.DAT.*
19: /SID2/gcgdata/geneseq/geneseq-emb1/AA1997.DAT.*
20: /SID2/gcgdata/geneseq/geneseq-emb1/AA1998.DAT.*
21: /SID2/gcgdata/geneseq/geneseq-emb1/AA1999.DAT.*
22: /SID2/gcgdata/geneseq/geneseq-emb1/AA2000.DAT.*
23: /SID2/gcgdata/geneseq/geneseq-emb1/AA2001.DAT.*
24: /SID2/gcgdata/geneseq/geneseq-emb1/AA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	ID	Description
1	1956	100.0	360 21 AAB09823	Endoglucanase prot
2	1956	100.0	360 23 AAO15054	Rhizopus arrhizus
3	1956	100.0	360 23 ABB08062	R. oryzae CP96001
4	1404	71.8	338 21 AAB09821	Endoglucanase prot
5	1404	71.8	338 23 AAO15052	Rhizopus arrhizus
6	1404	71.8	338 23 ABB08060	R. oryzae CP96001
7	1332	68.1	366 21 AAB09822	Endoglucanase prot
8	1332	68.1	366 23 AAO15053	Rhizopus arrhizus
9	1332	68.1	366 23 ABB08061	R. oryzae CP96001
10	1209	61.8	338 21 AAB09824	Endoglucanase prot

11	1209	61.8	338 23 AAO15055	Rhizopus arrhizus
12	1209	61.8	338 23 ABB08063	M. circinelloides
13	1195.5	61.1	387 21 AAB09825	Endoglucanase prot
14	1195.5	61.1	387 23 AAO15056	Rhizopus arrhizus
15	1195.5	61.1	387 23 ABB08064	M. circinelloides
16	1194	61.0	346 21 AAB09826	Endoglucanase prot
17	1194	61.0	346 23 AAO15057	Phycomyces nitens
18	1194	61.0	346 23 ABB08065	P. nitens CP99002
19	1067.5	54.6	245 23 AAO15063	Endoglucanase-rela
20	1005.5	51.4	228 23 AAO15062	Endoglucanase-rela
21	754	38.5	225 21 AAB04798	Amino acid sequenc
22	754	38.5	225 23 ABB05057	Chrysosporium Cl 8
23	747	38.2	225 17 AAB04935	Cellulytic enzyme
24	747	38.2	297 17 AAO04933	Chimeric endogluc
25	747	38.2	308 17 AAO04934	Cellulytic enzyme
26	745	38.1	299 17 AAB04928	Monocomponent endo
27	745	38.1	299 19 AAB03624	Thielavia terrestr
28	740	37.8	200 19 AAB03979	Hybrid DNA protein
29	736	37.6	306 19 AAB04270	Myceliophthora che
30	735	37.6	204 19 AAB03970	Thielavia terrestr
31	732	37.4	200 19 AAB03968	Humicola insolens
32	731.5	37.4	223 23 AAO15070	Humicola insolens
33	731.5	37.4	223 23 AAB08062	Sordaria fimicola
34	724	37.0	200 19 AAB03967	Fusarium oxysporum
35	722.5	36.9	376 12 AAB05272	Fusarium oxysporum
36	722.5	36.9	376 13 AAB05273	Endoglucanase #2
37	722.5	36.9	376 13 AAB05466	Cellulase containe
38	722.5	36.9	376 13 AAB05429	Endoglucanase enzy
39	722.5	36.9	376 13 AAB07969	Endoglucanase enzy
40	722.5	36.9	376 14 AAB02064	F. oxysporum endog
41	722.5	36.9	376 16 AAB07389	Fusarium oxysporum
42	722.5	36.9	376 19 AAB06617	Dye transfer inhib
43	716.5	36.6	376 14 AAB07151	Fusarium oxysporum
44	713	36.5	202 19 AAB03969	Fusarium oxysporum
45	712	36.4	357 15 ABB04127	Fusarium oxysporum

ALIGNMENTS

RESULT 1
AAB09823 standard; Protein; 360 AA.
XX
AC AAB09823;
XX
DT 25-SEP-2000 (first entry)
XX
DE Endoglucanase protein sequence 3.
XX
KW Endoglucanase; cellulose breakdown; produce pulp; papermaking;
KW animal foodstuffs;
XX
OS Rhizopus oryzae
XX
PN WO200024879-A1.
XX
PD 04-MAY-2000.
XX
PF 25-OCT-1999; 99WO-JP05884.
XX
PR 23-OCT-1998; 98JP-0302387.
XX
PA (MEIJ) MEIJU SEIKA KAISHA LTD.
XX
PI Nakamura Y, Moriya T, Baba Y, Yanai K, Sumida N, Nishimura T;
XX
PI Murashima K, Nakane A, Yaguchi T, Koga J, Murekami T, Kono T;
XX
DR WPI: 2000-365117/31.
XX
DR N-PSDB; AAB62728.
XX
PT Endoglucanases of fungal origin with high activity under alkaline
XX
PT conditions for production of paper pulp and animal feedstuffs -

PD 16-MAY-2002.
XX 12-NOV-2001; 2001WO-JP09858.
XX 10-NOV-2000; 2000JP-0343921.
XX (MEIJ) MEIJI SEIKA KAISHA LTD.
XX Koga J, Nakane A, Baba Y, Kono T;
XX WPI; 2002-471555/50.
DR Cellulase preparations containing transconjugant-originated
PT endoglucanase and non-ionic surfactants, useful in detergent
PT compositions, in treating cellulose fibers and delinking waste paper and
PT improving freeness of paper pulp -
XX Claim 3; Page 25-27; 38pp; Japanese.
XX The invention relates to a cellulase preparation comprising a
CC transconjugant-originated endoglucanase and a non-ionic surfactant. The
CC endoglucanase is selected from RCEI, RCEII, RCEIII, MCEI, MCEII or PCEI
CC proteins. The preparations are useful in detergent compositions, in
CC treating cellulose fibers and delinking waste paper and improving the
CC freeness of paper pulp. The fibers treated by the preparations have
CC reduced feathering and improved skin-feel and appearance with colour
CC clarification, local change in colour and softening, and after delinking
CC and paper pulp treatment, there is an improvement in freeness of the
CC paper pulp. This treatment with the cellulase preparation can be operated
CC at significantly lower cost. The present sequence represents the
CC R. oryzae CP960d1 RCEIII protein.
XX
SQ Sequence 360 AA;
Query Match 100.0%; Score 1956; DB 23; Length 360;
Best Local Similarity 100.0%; Pred. No. 2.6e-130;
Matches 360; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MKPFTTSSALALAVGTEMAHAECSEKAYQCCGKMDGPTCCSGSTCYDYPDNPPYS 60
DB 1 MKPFTTSSALALAVGTEMAHAECSEKAYQCCGKMDGPTCCSGSTCYDYPDNPPYS 60
QY 61 QCVNENLSTNKSHTKTTTESAKTTTGGSKTTTTEASKTTTTEASKTTTTEAS 120
DB 61 QCVNENLSTNKSHTKTTTESAKTTTGGSKTTTTEASKTTTTEASKTTTTEAS 120
QY 121 KKTITTTKASTSTSSSSASSTVNSAVSGASNGETTRWDCCKPSCWPGKADVTSP 180
DB 121 KKTITTTKASTSTSSSSASSTVNSAVSGASNGETTRWDCCKPSCWPGKADVTSP 180
QY 181 VGSCKNGKGTLDANNONGCVGSSSTYTCNDNQPVVSDLAAYFPAAASISGSEBATTWCCA 240
DB 181 VGSCKNGKGTLDANNONGCVGSSSTYTCNDNQPVVSDLAAYFPAAASISGSEBATTWCCA 240
QY 241 CFEFLTSTAVKGMVQVNTGSDLSNTGAHFDLQMPGGGVGVIYNGCATOMGAPTDG 300
DB 241 CFEFLTSTAVKGMVQVNTGSDLSNTGAHFDLQMPGGGVGVIYNGCATOMGAPTDG 300
QY 301 WGARVGVSSASDCSNLPSALQAGCKWRFGWFKADNPMTYKQVTCPKAITAKSGCSR 360
DB 301 WGARVGVSSASDCSNLPSALQAGCKWRFGWFKADNPMTYKQVTCPKAITAKSGCSR 360
RESULT 4
AAB09821
ID AAB09821 standard; Protein; 338 AA.
XX AAB09821;
XX 25-SEP-2000 (first entry)
XX Endoglucanase protein sequence 1.
XX

KW Endoglucanase; cellulose breakdown; produce pulp; papermaking;
KM animal feedstuff.
XX Rhizopus oryzae.
XX WO200024879-A1.
XX 04-MAY-2000.
XX 25-OCT-1999; 99WO-JP05864.
XX 23-OCT-1998; 98JP-0302387.
XX (MEIJ) MEIJI SEIKA KAISHA LTD.
XX Nakamura Y, Moriya T, Baba Y, Yanai K, Sumida N, Nishimura T;
XX Murashima K, Nakane A, Yaguchi T, Koga J, Murakami T, Kono T;
XX WPI; 2000-365117/31.
DR N-PSDB; AAA62726.
XX
PT Endoglucanases of fungal origin with high activity under alkaline
PT conditions for production of paper pulp and animal feedstuffs -
XX Claim 44; Page 106-108; 180pp; Japanese.
XX This sequence represents an endoglucanase protein. The invention relates
CC to an endoglucanase of fungal origin which can completely break down
CC purified cellulose at a concentration of less than 1mg protein/litre,
CC and produces more than 50% breakdown of cellulose at pH 8.5. The
CC invention includes endoglucanase protein sequences (see
CC AAB09825-B09830), endoglucanase nucleotide sequences (see
CC AAA62726-A62732), and primers (AAA62733-A62802) which are used in the
CC identification of the endoglucanase sequences, and in the construction of
CC vectors containing the polynucleotides. The endoglucanase enzymes are
CC used for the production of pulp for papermaking and for the production of
CC animal feedstuffs.
XX
SQ Sequence 338 AA;
Query Match 71.8%; Score 1404; DB 21; Length 338;
Best Local Similarity 73.7%; Pred. No. 2.3e-91;
Matches 266; Conservative 32; Mismatches 39; Indels 24; Gaps 7;
QY 1 MKPFTTSSALALAVGTEMAHAECSEKAYQCCGKMDGPTCCSGSTCYDYPDNPPYS 60
DB 1 MKPFTTSSALALAVGTEMAHAECSEKAYQCCGKMDGPTCCSGSTCYDYPDNPPYS 60
QY 61 QCVNENLSTNKSHTKTTTESAKTTTGGSKTTTTEASKTTTTEASKTTTTEAS 120
DB 61 QCVNENLSTNKSHTKTTTESAKTTTGGSKTTTTEASKTTTTEASKTTTTEAS 120
QY 59 QCLPSG--SSGNKSS-----ESAHKTTTAAHKTTTA-----AHKTTTAPAK 100
DB 59 QCLPSG--SSGNKSS-----ESAHKTTTAAHKTTTA-----AHKTTTAPAK 100
QY 121 KKTITTTKASTSTSSSSASSTVNSAVSGASNGETTRWDCCKPSCWPGKADVTSP 180
DB 121 KKTITTTKASTSTSSSSASSTVNSAVSGASNGETTRWDCCKPSCWPGKADVTSP 180
QY 101 K-TTVAKAST-PSNSSSSSGKXSAVSGASNGVYTRWDCCKKASCWPGKANVSSP 157
DB 101 K-TTVAKAST-PSNSSSSSGKXSAVSGASNGVYTRWDCCKKASCWPGKANVSSP 157
QY 181 VGSCKNGKGTLDANNONGCVGSSSTYTCNDNQPVVSDLAAYFPAAASISGSEBATTWCCA 240
DB 181 VGSCKNGKGTLDANNONGCVGSSSTYTCNDNQPVVSDLAAYFPAAASISGSEBATTWCCA 240
QY 240 ACFELTSTAVKGMVQVNTGSDLSNTGAHFDLQMPGGGVGVIYNGCATOMGAPTD 299
DB 240 ACFELTSTAVKGMVQVNTGSDLSNTGAHFDLQMPGGGVGVIYNGCATOMGAPTD 299
QY 218 SCFELTSTAVKGMVQVNTGSDLSNTGAHFDLQMPGGGVGVIYNGCATOMGAPTD 277
DB 218 SCFELTSTAVKGMVQVNTGSDLSNTGAHFDLQMPGGGVGVIYNGCATOMGAPTD 277
QY 300 WGARVGVSSASDCSNLPSALQAGCKWRFGWFKADNPMTYKQVTCPKAITAKSGCSR 359
DB 300 WGARVGVSSASDCSNLPSALQAGCKWRFGWFKADNPMTYKQVTCPKAITAKSGCSR 359
QY 278 WGSRYGIGSSASDCSNLPSALQAGCKWRFGWFKADNPMTYKQVTCPKAITAKSGCSR 337
DB 278 WGSRYGIGSSASDCSNLPSALQAGCKWRFGWFKADNPMTYKQVTCPKAITAKSGCSR 337
QY 360 K 360
DB 338 K 338

CC binding domain. The zygomycetes-originated endoglucanase enzymes of the
 CC invention have enhanced endoglucanase activity. The zygomycetes-
 CC originated endoglucanase enzymes of the invention are useful for
 CC processing fibres, de-inking waste paper and improving the freeness of
 CC paper pulp - which is particularly applicable in detergent compositions.
 CC The present amino acid sequence represents an endoglucanase-related
 CC protein of the invention.

XX Sequence 366 AA;

Query Match 68.1%; Score 1332; DB 23; Length 366;

Best Local Similarity 66.4%; Pred. No. 3.1e-86; Indels 24; Gaps 5;

Matches 249; Conservative 39; Mismatches 63;

DB 1 MKFLITASSALILAVGTEMAHAEGSKAYOCCGKMDGPTCCESGTCVDYDPNPFYS 60
 1 MKFLITSSALILAVGTEMAHAEGSKAYOCCGKMDGPTCCESGTC--KVANDYYS 58
 QY 61 QCVNENILSTNKS-----HKTTSAAKTTTKSKKTTTEASKTTT 106
 DB 59 QCLAPE--SNKNSSSECKLYGCGGKMDGPTCCESGTCVKVSNDYYSQCLAPESNGK 116
 QY 107 TTEASKTTTTEASKTTTTKKASTSTSSSSASTNYSAVSGASNGETTRYWDCK 166
 DB 117 TSESAAKTTTTPAKKITTATAKSNSSSG-----KSIYSGASNGVTRRYWDCK 171
 QY 167 PSCSWPGKADVTSPVSGCNKDKGT-LADNNTONGCVGSSSYTNDNQPVVSDILAYGFA 225
 DB 172 ASCSWPGKANVSVPKSCNKGVTALSDSNVSGCNGNSYMCNDNQPVAVNDNLAYGFA 231
 QY 226 AASISGSEATWCCAFELTFTSTAVYGGKRVVNTGSDLSNTGAHFDDLOMPGGVG 285
 DB 232 AASISGSESEWCCSCEFLTFTSTVAGKRVIVNTGDLGSSSTAHDLOMPGGVG 291
 QY 286 IYNGCATOWGAPTDGMDGARYGVSSASDCSNLPSALQAGCKMRPFKADNPTTYKQV 345
 DB 292 IFNGSKQWGANPDGMSRYGSISSADCSLPSALQAGCKMRPFKADNPTTYKEY 351
 QY 346 TCPKAITAKSGCSRK 360
 DB 352 TCPKEITAKTGCSRK 366

RESULT 9
 ABB08061
 ID ABB08061 standard; protein; 366 AA.

XX ABB08061;

DT 27-AUG-2002 (first entry)

DE R. oryzae CP96001 RCEII protein.

XX Cellulase; endoglucanase; surfactant; detergent; cellulose; paper;
 KM pulp treatment; RCEII.

XX Rhizopus oryzae.

XX Key Location/Qualifiers

FT Peptide 1..23

FT Protein /note= "signal peptide"

FT /note= "mature protein"

XX WO200238754-A1.

XX 16-MAY-2002.

XX 12-NOV-2001; 2001WO-JP09858.

XX 10-NOV-2000; 2000JP-0343921.

XX (MEIJ) MEIJI SEIKA KAISHA LTD.

XX Koga J, Nakane A, Baba Y, Kono T;
 PI WPI; 2002-471555/50.
 DR Cellulase preparations containing transconjugant-originated
 XX cellulase and non-ionic surfactants, useful in detergent
 PT compositions, in treating cellulose fibers and deinking waste paper and
 PT improving freeness of paper pulp

XX Claim 3; Page 23-24; 38pp; Japanese.

CC The invention relates to a cellulase preparation comprising a
 CC transconjugant-originated endoglucanase and a non-ionic surfactant. The
 CC endoglucanase is selected from RCEI, RCEII, RCEIII, MCEII or PCBI
 CC proteins. The preparations are useful in detergent compositions, in
 CC treating cellulose fibers and deinking waste paper and improving the
 CC freeness of paper pulp. The fibers treated by the preparations have
 CC reduced feathering and improved skin-feel and appearance with colour
 CC clarification, local change in colour and softening, and after deinking
 CC paper pulp. This treatment with the cellulase preparation can be operated
 CC at significantly lower cost. The present sequence represents the
 CC R. oryzae CP96001 RCEII protein.

SQ Sequence 366 AA;

Query Match 68.1%; Score 1332; DB 23; Length 366;

Best Local Similarity 66.4%; Pred. No. 3.1e-86; Indels 24; Gaps 5;

Matches 249; Conservative 39; Mismatches 63;

DB 1 MKFLITASSALILAVGTEMAHAEGSKAYOCCGKMDGPTCCESGTCVDYDPNPFYS 60
 1 MKFLITSSALILAVGTEMAHAEGSKAYOCCGKMDGPTCCESGTC--KVANDYYS 58
 QY 61 QCVNENILSTNKS-----HKTTSAAKTTTKSKKTTTTEASKTTT 106
 DB 59 QCLAPE--SNKNSSSECKLYGCGGKMDGPTCCESGTCVKVSNDYYSQCLAPESNGK 116
 QY 107 TTEASKTTTTEASKTTTTKKASTSTSSSSASTNYSAVSGASNGETTRYWDCK 166
 DB 117 TSESAAKTTTTPAKKITTATAKSNSSSG-----KSIYSGASNGVTRRYWDCK 171
 QY 167 PSCSWPGKADVTSPVSGCNKDKGT-LADNNTONGCVGSSSYTNDNQPVVSDILAYGFA 225
 DB 172 ASCSWPGKANVSVPKSCNKGVTALSDSNVSGCNGNSYMCNDNQPVAVNDNLAYGFA 231
 QY 226 AASISGSEATWCCAFELTFTSTAVYGGKRVVNTGSDLSNTGAHFDDLOMPGGVG 285
 DB 232 AASISGSESEWCCSCEFLTFTSTVAGKRVIVNTGDLGSSSTAHDLOMPGGVG 291
 QY 286 IYNGCATOWGAPTDGMDGARYGVSSASDCSNLPSALQAGCKMRPFKADNPTTYKQV 345
 DB 292 IFNGSKQWGANPDGMSRYGSISSADCSLPSALQAGCKMRPFKADNPTTYKEY 351
 QY 346 TCPKAITAKSGCSRK 360
 DB 352 TCPKEITAKTGCSRK 366

RESULT 10
 AAB09824
 ID AAB09824 standard; Protein; 338 AA.

XX AAB09824;

DT 25-SEP-2000 (first entry)

DE Endoglucanase protein sequence 4.

XX Endoglucanase; cellulose breakdown; produce pulp; papermaking;
 KW animal foodstuff.

OS Mucor circinelloides.
 XX WO200024879-A1.
 XX 04-MAY-2000.
 XX
 XX 25-OCT-1999; 99WO-JP05884.
 XX
 XX 23-OCT-1998; 98JP-0302387.
 XX
 XX (MEIJ) MEIJ SEIKA KAISHA LTD.
 XX Nakamura Y, Moriya T, Baba Y, Yanai K, Sumida H, Nishimura T;
 PI Murashima K, Nakane A, Yaguchi T, Koga J, Murakami T, Kono T;
 XX WPI, 2000-365117/31.
 DR N-PSDB; AAA62729.
 XX
 XX Endoglucanases of fungal origin with high activity under alkaline
 PT conditions for production of paper pulp and animal feedstuffs -
 PS Claim 44; Page 120-122; 180pp; Japanese.
 XX
 XX This sequence represents an endoglucanase protein. The invention relates
 CC to an endoglucanase of fungal origin which can completely break down
 CC purified cellulose at a concentration of less than 1mg protein/litre,
 CC and produces more than 50% breakdown of cellulose at pH 8.5. The
 CC invention includes endoglucanase protein sequences (see
 CC AB09825-B09830), endoglucanase nucleotide sequences (see
 CC AAA62726-A62732) and primers (AAA62733-A62802) which are used in the
 CC identification of the endoglucanase sequences, and in the construction of
 CC vectors containing the polynucleotides. The endoglucanase enzymes are
 CC used for the production of pulp for papermaking and for the production of
 CC animal feedstuffs.
 XX
 XX Sequence 338 AA;
 SQ
 Query Match 61.8%; Score 1209; DB 21; Length 338;
 Best Local Similarity 61.1%; Pred. No. 1.4e-77;
 Matches 220; Conservative 42; Mismatches 76; Indels 22; Gaps 5;
 QY 1 MKPFTIASIALALAVGTEMAHAABCSKAYVQCGKMDGPTCCSGSTCYVDYDNPYS 60
 DB 1 MKPFTVAITSIALVALASSS-ABAASCSGVGCGGIGMSGPTCCSGSTCVAQSGNKYYS 59
 QY 61 QCVNENLSTNKSHTTTTBSAKTTTSGSKTTTTEASKTTTTEASKTTTTEAS 120
 DB 60 QCLPGSHSNAGNANS-----STKKTST---KSTTTAKATATVTTKTKTT----- 103
 QY 121 KKTITTTKASTSTSSSSSSASTNYSAVSGASNGETTRVMDCKPSCSWPGKADYTS 180
 DB 104 --TKTTTSTTAALASTSTSSAGKVIISGKSGSGSTTRVMDCKKASCSPGKASVTGP 161
 QY 181 VGSCKNGKTLADNNTONGCVGSSSYTCNDNQPVVSDDLAYGFAAASISGSSSATWCCA 240
 DB 162 VDTCASNGISILDANAOSGCGNGFCMNNQNPAAVNDLAYGFAAASISGSSSATWCCA 221
 QY 241 CFEITFTSTAVKGMVVOVNTGSDLSNTGAHFDLMPGCGVIGYNGCATONGAPITDG 300
 DB 222 CYELTFTSGAASGKGMVVOVNTGSDLSN---HFDLMPGCGVIGYNGCAOAGAPITDG 278
 QY 301 WGARVGVSSASDCSNLPSALQAGCKWRFGMFKXNDNPTMTYKQVTCPEKAITTASGSRK 360
 DB 279 WGARVGVSSVSDCASLPSALQAGCKWRFGMFKXNDNPTMTYKQVTCPEKAITTASGSRK 338
 RESULT 11
 ID AAO15055 standard; Protein; 338 AA.
 XX AAO15055;
 XX
 DT 22-AUG-2002 (first entry)

XX Rhizopus arrhizus endoglucanase-related protein 4.
 DE
 XX Zygomycetes-originated endoglucanase; cellulose binding domain;
 KW fibre processing; waste paper de-inking; paper pulp.
 XX
 XX Mucor circinelloides.
 OS
 XX WO200242474-A1.
 XX 30-MAY-2002.
 XX
 XX 21-NOV-2001; 2001WO-JP10188.
 XX
 XX 21-NOV-2000; 2000JP-0354296.
 XX
 XX (MEIJ) MEIJ SEIKA KAISHA LTD.
 XX Nakane A, Baba Y, Koga J, Kubota H;
 PI WPI, 2002-471729/50.
 DR N-PSDB; AAA43247.
 XX
 XX Cellulose-binding domain-lacking Zygomycetes-originated endoglucanase,
 PT with effect of endoglucanase activity enhanced in processing fibers,
 PT deinking waste paper and improving freeness of paper pulp -
 PS Claim 5; Page 68-70; 109pp; Japanese.
 XX
 XX The invention comprises the amino acid and coding sequences of
 CC zygomycetes-originated endoglucanase enzymes lacking the cellulose
 CC binding domain. The zygomycetes-originated endoglucanase enzymes of the
 CC invention have enhanced endoglucanase activity. The zygomycetes-
 CC originated endoglucanase enzymes of the invention are useful for
 CC processing fibres, de-inking waste paper and improving the freeness of
 CC paper pulp - which is particularly applicable in detergent compositions.
 CC The present amino acid sequence represents an endoglucanase-related
 CC protein of the invention.
 XX
 XX Sequence 338 AA;
 SQ
 Query Match 61.8%; Score 1209; DB 23; Length 338;
 Best Local Similarity 61.1%; Pred. No. 1.4e-77;
 Matches 220; Conservative 42; Mismatches 76; Indels 22; Gaps 5;
 QY 1 MKPFTIASIALALAVGTEMAHAABCSKAYVQCGKMDGPTCCSGSTCYVDYDNPYS 60
 DB 1 MKPFTVAITSIALVALASSS-ABAASCSGVGCGGIGMSGPTCCSGSTCVAQSGNKYYS 59
 QY 61 QCVNENLSTNKSHTTTTBSAKTTTSGSKTTTTEASKTTTTEASKTTTTEAS 120
 DB 60 QCLPGSHSNAGNANS-----STKKTST---KSTTTAKATATVTTKTKTT----- 103
 QY 121 KKTITTTKASTSTSSSSSSASTNYSAVSGASNGETTRVMDCKPSCSWPGKADYTS 180
 DB 104 --TKTTTSTTAALASTSTSSAGKVIISGKSGSGSTTRVMDCKKASCSPGKASVTGP 161
 QY 181 VGSCKNGKTLADNNTONGCVGSSSYTCNDNQPVVSDDLAYGFAAASISGSSSATWCCA 240
 DB 162 VDTCASNGISILDANAOSGCGNGFCMNNQNPAAVNDLAYGFAAASISGSSSATWCCA 221
 QY 241 CFEITFTSTAVKGMVVOVNTGSDLSNTGAHFDLMPGCGVIGYNGCATONGAPITDG 300
 DB 222 CYELTFTSGAASGKGMVVOVNTGSDLSN---HFDLMPGCGVIGYNGCAOAGAPITDG 278
 QY 301 WGARVGVSSASDCSNLPSALQAGCKWRFGMFKXNDNPTMTYKQVTCPEKAITTASGSRK 360
 DB 279 WGARVGVSSVSDCASLPSALQAGCKWRFGMFKXNDNPTMTYKQVTCPEKAITTASGSRK 338
 RESULT 12
 ID ABB08063 standard; protein; 338 AA.
 XX ABB08063
 XX

XX AC ABB08063;
 XX DT 27-AUG-2002 (first entry)
 XX DE M. circinelloides CP99001 MCEI protein.
 XX DE Cellulase; endoglucanase; surfactant; detergent; cellulose; paper;
 XX KM pulp treatment; MCEI.
 XX OS Mucor circinelloides.
 XX FH Key Location/Qualifiers
 XX FT Peptide 1..22
 XX FT /note= "signal peptide"
 XX FT 23..338
 XX FT Protein /note= "mature protein"
 XX FT
 XX PN WO200238754-A1.
 XX PD 16-MAY-2002.
 XX PF 12-NOV-2001; 2001WO-JP09858.
 XX PR 10-NOV-2000; 2000JP-0343921.
 XX PA (MEIJ) MEIJI SEIKA KAISHA LTD.
 XX PI Koga J., Nakane A., Baba Y., Kono T;
 XX DR WPI; 2002-471555/50.
 XX XX
 XX PT Cellulase preparations containing transconjugant-originated
 XX PT endoglucanase and non-ionic surfactants, useful in detergent
 XX PT compositions, in treating cellulose fibers and delinking waste paper and
 XX PT improving freeness of paper pulp -
 XX PS Claim 3; Page 27-29; 38pp; Japanese.
 XX XX
 XX CC The invention relates to a cellulase preparation comprising a
 XX CC transconjugant-originated endoglucanase and a non-ionic surfactant. The
 XX CC endoglucanase is selected from RCEI, RCEII, MCEI, MCEII or PC-EI
 XX CC proteins. The preparations are useful in detergent compositions, in
 XX CC treating cellulose fibers and delinking waste paper and improving the
 XX CC freeness of paper pulp. The fibers treated by the preparations have
 XX CC reduced feathering and improved skin-feel and appearance with colour
 XX CC clarification, local change in colour and softening, and after delinking
 XX CC and paper pulp treatment, there is an improvement on freeness of the
 XX CC paper pulp. This treatment with the cellulase preparation can be operated
 XX CC at significantly lower cost. The present sequence represents the
 XX CC M. circinelloides CP99001 MCEI protein.
 XX SQ Sequence 338 AA;
 XX
 XX Query Match 61.8%; Score 1209; DB 23; Length 338;
 XX Best Local Similarity 61.1%; Pred. No. 1.4e-77;
 XX Matches 220; Conservative 42; Mismatches 76; Indels 22; Gaps 5;

OY 241 CPELTSTAVKGMVYQVNTSGSDLSNTGAHFIDQMPGGVGIYNGCATOWGAPTDG 300
 DB 222 CYELFTSGASGKMMVQVNTSGDLSN---HFDLQMPGGVGIYNGCAOWGABNDG 278
 OY 301 WGARVGVSSASDCSNLPSALQAGCKMRFKXNDPTMYKQVTPKXITAKSGSRK 360
 DB 279 WGARVGVSSVSDCASLPSALQAGCKMRFKXNDPTMYKQVTPKXITAKSGSRK 338
 RESULT 13
 AAB09825
 ID AAB09825 standard; Protein, 387 AA.
 XX
 XX AC AAB09825;
 XX DT 25-SEP-2000 (first entry)
 XX DE Endoglucanase protein sequence 5.
 XX DE Endoglucanase; cellulose breakdown; produce pulp; papermaking;
 XX KM animal feedstuff.
 XX OS Phycomyces nitens.
 XX PN WO200024879-A1.
 XX PD 04-MAY-2000.
 XX PF 25-OCT-1999; 99WO-JP05884.
 XX PR 23-OCT-1998; 98JP-0302387.
 XX PA (MEIJ) MEIJI SEIKA KAISHA LTD.
 XX PI Nakamura Y., Moriya T., Baba Y., Yanai K., Sumida N., Nishimura T;
 XX PI Mureshima K., Nakane A., Yaguchi T., Koga J., Murakami T., Kono T;
 XX DR WPI; 2000-365117/31.
 XX DR N-PSDB; AAA62730.
 XX PT Endoglucanases of fungal origin with high activity under alkaline
 XX PT conditions for production of paper pulp and animal feedstuffs -
 XX PS Claim 44; Page 125-127; 180pp; Japanese.
 XX XX
 XX CC This sequence represents an endoglucanase protein. The invention relates
 XX CC to an endoglucanase of fungal origin which can completely break down
 XX CC purified cellulose at a concentration of less than 1mg protein/litre,
 XX CC and produces more than 50% breakdown of cellulose at pH 8.5. The
 XX CC invention includes endoglucanase protein sequences (see
 XX CC AAB09825-B09830), endoglucanase nucleotide sequences (see
 XX CC AAA62726-A62732) and primers (AAA62733-A62802) which are used in the
 XX CC identification of the endoglucanase sequences, and in the construction of
 XX CC vectors containing the polynucleotides. The endoglucanase enzymes are
 XX CC used for the production of pulp for papermaking and for the production of
 XX CC animal feedstuffs.
 XX SQ Sequence 387 AA;
 XX
 XX Query Match 61.1%; Score 1195.5; DB 21; Length 387;
 XX Best Local Similarity 56.2%; Pred. No. 1.4e-76;
 XX Matches 221; Conservative 47; Mismatches 86; Indels 39; Gaps 6;

```

Db 120 GNASTKKTSTKTS--TTAKATATVTTKVTKTTTSTTAASSTSSAGYKV 177
Qy 148 VSGASGNGETTRWDCCKSPGKADVTSPVSGCNKDKTLADNNTONGCVGSSYT 207
Db 178 ISGKSGSGSTTRWDCCKASCWPGRKASVTGPDVTCASNGISILDANAOSGCGNGFM 237
Qy 208 CNDQPMVWSDLLAYGPAASISGGSSEATWCACFELTFTSTAVKGRKVVQVNTGSDL 267
Db 238 CNDQPMVWSDLLAYGPAASISGGSSEATWCACFELTFTSTAVKGRKVVQVNTGSDL 297
Qy 268 GSNTHAFDLOMPGSGVGIYNGCATOWGAPTDGNGARYGVSSASDCSNLPSALOAGCKW 327
Db 298 GSN--HFDLOMPGSGVGIYNGCATOWGAPTDGNGARYGVSSASDCSNLPSALOAGCKW 354
Qy 328 RFGWFKNADNPTMTYKQVTCPKAITAKSGCSRK 360
Db 355 RFWNFKNSDNPMTMFKVTCPELITRSGCERK 387

```

RESULT 14

ID AAO15056 standard; Protein; 387 AA.

AAO15056;

22-AUG-2002 (first entry)

Rhizopus arrhizus endoglucanase-related protein 5.

Zygomycetes-originated endoglucanase; cellulose binding domain;

fibres processing; waste paper de-inking; paper pulp.

Mucor circinelloides.

MO200242474-A1.

30-MAY-2002.

21-NOV-2001; 2001WO-JP10188.

21-NOV-2000; 2000JP-0354296.

(MEIJ) MEIJI SEIKA KAISHA LTD.

Nakane A, Baba Y, Koga J, Kubota H;

WPI; 2002-471729/50.

N-PSDB; AAL43248.

Cellulose-binding domain-lacking Zygomycetes-originated endoglucanase,

with effect of endoglucanase activity enhanced in processing fibers,

deinking waste paper and improving freeness of paper pulp -

Claim 5; Page 73-75; 109pp; Japanese.

The invention comprises the amino acid and coding sequences of
zygomycetes-originated endoglucanase enzymes lacking the cellulose
binding domain. The zygomycetes-originated endoglucanase enzymes of the
invention have enhanced endoglucanase activity. The zygomycetes-
originated endoglucanase enzymes of the invention are useful for
processing fibres, de-inking waste paper and improving the freeness of
paper pulp - which is particularly applicable in detergent compositions.
The present amino acid sequence represents an endoglucanase-related
protein of the invention.

Sequence 387 AA;

Query Match / 61.1%; Score 1195.5; DB 23; length 387;

Best Local Similarity 56.2%; Pred. No. 1.4e-76;

Matches 221; Conservative 47; Mismatches 86; Indels 39; Gaps 6;

Qy 1 MKPFIITASSALIALAVGTEMMAAECRAVYQCGKMDGPTCCSGSGSTCVDPDNPFRYS 60

```

Db 1 MKFTVATISIAVALALSSS-ABAAACSSVYGQCGGIGMTGPTCCDAGSTCAQKDNKYS 59
Qy 61 QCPVNEULTSTNKSHT-----TTESAKTTTIGSK----- 94
Db 60 QCPVNEULTSTNKSHT-----TTESAKTTTIGSK----- 119
Qy 95 -----TTTTEASKTTTTEASKTTT-TTTKAISTSSSSASTVYSA 147
Db 120 GNASTKKTSTKTS--TTAKATATVTTKVTKTTTSTTAASSTSSAGYKV 177
Qy 148 VSGASGNGETTRWDCCKSPGKADVTSPVSGCNKDKTLADNNTONGCVGSSYT 207
Db 178 ISGKSGSGSTTRWDCCKASCWPGRKASVTGPDVTCASNGISILDANAOSGCGNGFM 237
Qy 208 CNDQPMVWSDLLAYGPAASISGGSSEATWCACFELTFTSTAVKGRKVVQVNTGSDL 267
Db 238 CNDQPMVWSDLLAYGPAASISGGSSEATWCACFELTFTSTAVKGRKVVQVNTGSDL 297
Qy 268 GSNTHAFDLOMPGSGVGIYNGCATOWGAPTDGNGARYGVSSASDCSNLPSALOAGCKW 327
Db 298 GSN--HFDLOMPGSGVGIYNGCATOWGAPTDGNGARYGVSSASDCSNLPSALOAGCKW 354
Qy 328 RFGWFKNADNPTMTYKQVTCPKAITAKSGCSRK 360
Db 355 RFWNFKNSDNPMTMFKVTCPELITRSGCERK 387

```

RESULT 15

ID ABB08064 standard; protein; 387 AA.

ABB08064;

27-AUG-2002 (first entry)

M. circinelloides CP99001 MCEII protein.

Cellulase; endoglucanase; surfactant; detergent; cellulose; paper;

pulp treatment; MCEII.

Mucor circinelloides.

Key Location/Qualifiers

Peptide 1..22 /note= "signal peptide"

Protein 23..387 /note= "mature protein"

MO200238754-A1.

16-MAY-2002.

12-NOV-2001; 2001WO-JP09858.

10-NOV-2000; 2000JP-0343921.

(MEIJ) MEIJI SEIKA KAISHA LTD.

Koga J, Nakane A, Baba Y, Kono T;

WPI; 2002-471555/50.

Cellulase preparations containing transconjugant-originated
endoglucanase and non-ionic surfactants, useful in detergent
compositions, in treating cellulose fibers and deinking waste paper and
improving freeness of paper pulp -

Claim 3; Page 29-31; 38pp; Japanese.

The invention relates to a cellulase preparation comprising a
transconjugant-originated endoglucanase and a non-ionic surfactant. The
endoglucanase is selected from RCEI, RCEII, MCEII, MCEII or PCEI

CC proteins. The preparations are useful in detergent compositions, in
CC treating cellulose fibers and delinking waste paper and improving the
CC freeness of paper pulp. The fibers treated by the preparations have
CC reduced feathering and improved skin-feel and appearance with colour
CC clarification, local change in colour and softening, and after delinking
CC and paper pulp treatment, there is an improvement on freeness of the
CC paper pulp. This treatment with the cellulase preparation can be operated
CC at significantly lower cost. The present sequence represents the
CC M. circinelloides CP99001 MCEII protein.

XX
SQ Sequence 387 AA;

Query Match 61.1%; Score 1195.5; DB 23; Length 387;
Best Local Similarity 56.2%; Pred. No. 1.4e-76;
Matches 221; Conservative 47; Mismatches 86; Indels 39; Gaps 6;

1 MKFLTIASSAIALAVGTEMAHAECSEKAYOCGKNDGPTCCSGSTCVDPNPFPYS 60
1 MKFTVAITSIYVALALSS-AEASCSYVGCGGIGMTGPTCCDAGSTCAQKDKNKYYS 59
61 QCVNENLTSTNKSHT-----TTESAKTTTKGSK-----94
60 QCIPEKSGSSSSSCSVYQCGIGMSPTCCSGSTCVAQEGNKYEQCLPGSHSNA 119
95 -----KTTTEASKTTTTEASKTTTTEASKTTT-TTKKASTSTSSSSASTNYS 147
120 GNASTKKTSTKTS--TTAKATATVTTKTIVTKTKTKTKTSTTAASTSTSSAGYK 177
148 VSGASNGETTRWDCCKSPGKADVTSPVSGCNKDKTLADNNTONGCVGGSYT 207
178 ISGKSGSGSTTRWDCCKASCPGASVTGPVDTCASNGISLDANAGCGNGNGFM 237
208 CNDNQPWVSDDLAYGFPAASISGSEATWCCACFELTFTSTAVKSKMVOYTNNGSD 267
238 CNNNPMAVNDLAYGFPAASISGSEATWCCACFELTFTSTAVKSKMVOYTNNGSD 297
268 GSNTGAHFDLQMPGGGVCITNGCATOWGAPTDGNGARYGVSSASDCSNLPSALQAGCKW 327
298 GSN--HFDLQMPGGGVCITNGCATOWGAPTDGNGARYGVSSASDCSNLPSALQAGCKW 354
328 RFGWFKADNPMTYKQVTCPRATYAKSGCSRK 360
355 RFWFKNSDNPMTFKEVTCPRATYAKSGCSRK 387

Search completed: June 18, 2003, 15:30:53
Job time : 40.4567 secs